



## SEQUENCE LISTING

&lt;110&gt; BORRELLI, MICHAEL J.

<120> METHODS AND COMPOSITIONS FOR HEAT ACTIVATED GENE  
THERAPY USING CYTOLETHAL DISTENDING TOXIN

&lt;130&gt; 10546-109

&lt;140&gt; 10/764,316

&lt;141&gt; 2004-01-23

&lt;150&gt; 60/442,473

&lt;151&gt; 2003-01-24

&lt;160&gt; 25

&lt;170&gt; PatentIn Ver. 3.3

&lt;210&gt; 1

&lt;211&gt; 2384

&lt;212&gt; DNA

&lt;213&gt; Haemophilus ducreyi

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&lt;213&gt; Campylobacter jejuni

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 gagaaaacac gataatgaag gttctggaag caaaaaggccc aaggtggagt attcagaaga 2340  
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 gagtctaccc gacataagtc gagggacttt atgtttttga ggctttctgt tgccatggtg 2640  
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<210> 7  
 <211> 2493  
 <212> DNA  
 <213> Homo sapiens

<400> 7  
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 gcggggtcgg ggagggtgcaa aaggatgaaa agcccgtgga agcggagctg agcagatccg 120  
 agccgggctg gcggcagaga aaccgcaggg agagcctcac tgctgagcgc ccctcgacgg 180  
 cggagcggca gcagcctccg tggcctccag catccgacaa gaagcttcag ccatgcaggc 240  
 cccacgggag ctgcggtgg gcatcgacct gggcaccacc tactcgtgcg tgggctgttt 300  
 tcagcagggc cgcgtggaga tcctggccaa cgaccagggc aaccgcacca cgcccagcta 360  
 cgtggccttc accgacaccg agcggctggt cggggacgcy gccaaagagcc aggcggccct 420  
 gaacccccac aacaccgtgt tcgatgcaa gcggtgatc gggcgcaagt tcgaggacac 480  
 cacggtgcag tcggacatga agcactggcc cttccagggtg gtgagcgagg gcggcaagcc 540  
 caagggtgcy gtatgctacc ggggggagga caagacgttc taccgccagg agatctcgtc 600  
 catggtgctg agcaagatga aggagacggc cgaggcgtac ctgggccagc ccgtgaagca 660  
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 cactgctgga gataccacc tgggaggaga ggacttcgac aaccggctcg tgaaccactt 960  
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 ccaccaagca gaccagact ttcaccacct actcggacaa ccagcctggg gtcttcatcc 1560  
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 acaaccagct ggcagagaag gaggagtatg agcatcagaa gaggggagctg gagcaaatct 2040  
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 gcgctcaagc ccaccagggg gaccccagca cgggccccat cattgaggag gttgatgaa 2160  
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 ctttcaccta tattttgtgt attttgttac ttgtatgtat gaattttgtt atgtaaaata 2460  
 tagttataga cctaaataaa cttttaaaac tcc 2493

<210> 8  
 <211> 643  
 <212> PRT  
 <213> Homo sapiens

<400> 8  
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 1 5 10 15

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			20					25					30				
Asn	Asp	Gln	Gly	Asn	Arg	Thr	Thr	Pro	Ser	Tyr	Val	Ala	Phe	Thr	Asp		
		35					40					45					
Thr	Glu	Arg	Leu	Val	Gly	Asp	Ala	Ala	Lys	Ser	Gln	Ala	Ala	Leu	Asn		
	50					55					60						
Pro	His	Asn	Thr	Val	Phe	Asp	Ala	Lys	Arg	Leu	Ile	Gly	Arg	Lys	Phe		
	65				70					75					80		
Ala	Asp	Thr	Thr	Val	Gln	Ser	Asp	Met	Lys	His	Trp	Pro	Phe	Arg	Val		
				85					90					95			
Val	Ser	Glu	Gly	Gly	Lys	Pro	Lys	Val	Arg	Val	Cys	Tyr	Arg	Gly	Glu		
			100					105					110				
Asp	Lys	Thr	Phe	Tyr	Pro	Glu	Glu	Ile	Ser	Ser	Met	Val	Leu	Ser	Lys		
		115					120					125					
Met	Lys	Glu	Thr	Ala	Glu	Ala	Tyr	Leu	Gly	Gln	Pro	Val	Lys	His	Ala		
	130					135					140						
Val	Ile	Thr	Val	Pro	Ala	Tyr	Phe	Asn	Asp	Ser	Gln	Arg	Gln	Ala	Thr		
	145				150					155					160		
Lys	Asp	Ala	Gly	Ala	Ile	Ala	Gly	Leu	Asn	Val	Leu	Arg	Ile	Ile	Asn		
			165					170						175			
Glu	Pro	Thr	Ala	Ala	Ala	Ile	Ala	Tyr	Gly	Leu	Asp	Arg	Arg	Gly	Ala		
			180					185					190				
Gly	Glu	Arg	Asn	Val	Leu	Ile	Phe	Asp	Leu	Gly	Gly	Gly	Thr	Phe	Asp		
		195					200					205					
Val	Ser	Val	Leu	Ser	Ile	Asp	Ala	Gly	Val	Phe	Glu	Val	Lys	Ala	Thr		
	210					215					220						
Ala	Gly	Asp	Thr	His	Leu	Gly	Gly	Glu	Asp	Phe	Asp	Asn	Arg	Leu	Val		
	225				230					235					240		
Asn	His	Phe	Met	Glu	Glu	Phe	Arg	Arg	Lys	His	Gly	Lys	Asp	Leu	Ser		
			245						250					255			
Gly	Asn	Lys	Arg	Ala	Leu	Arg	Arg	Leu	Arg	Thr	Ala	Cys	Glu	Arg	Ala		
			260					265					270				
Lys	Arg	Thr	Leu	Ser	Ser	Ser	Thr	Gln	Ala	Thr	Leu	Glu	Ile	Asp	Ser		
		275					280					285					
Leu	Phe	Glu	Gly	Val	Asp	Phe	Tyr	Thr	Ser	Ile	Thr	Arg	Ala	Arg	Phe		
	290					295					300						
Glu	Glu	Leu	Cys	Ser	Asp	Leu	Phe	Arg	Ser	Thr	Leu	Glu	Pro	Val	Glu		
	305				310					315					320		

Lys Ala Leu Arg Asp Ala Lys Leu Asp Lys Ala Gln Ile His Asp Val  
 325 330 335  
 Val Leu Val Gly Gly Ser Thr Arg Ile Pro Lys Val Gln Lys Leu Leu  
 340 345 350  
 Gln Asp Phe Phe Asn Gly Lys Glu Leu Asn Lys Ser Ile Asn Pro Asp  
 355 360 365  
 Glu Ala Val Ala Tyr Gly Ala Ala Val Gln Ala Ala Val Leu Met Gly  
 370 375 380  
 Asp Lys Cys Glu Lys Val Gln Asp Leu Leu Leu Leu Asp Val Ala Pro  
 385 390 395 400  
 Leu Ser Leu Gly Leu Glu Thr Ala Gly Gly Val Met Thr Thr Leu Ile  
 405 410 415  
 Gln Arg Asn Ala Thr Ile Pro Thr Lys Gln Thr Gln Thr Phe Thr Thr  
 420 425 430  
 Tyr Ser Asp Asn Gln Pro Gly Val Phe Ile Gln Val Tyr Glu Gly Glu  
 435 440 445  
 Arg Ala Met Thr Lys Asp Asn Asn Leu Leu Gly Arg Phe Glu Leu Ser  
 450 455 460  
 Gly Ile Pro Pro Ala Pro Arg Gly Val Pro Gln Ile Glu Val Thr Phe  
 465 470 475 480  
 Asp Ile Asp Ala Asn Gly Ile Leu Ser Val Thr Ala Thr Asp Arg Ser  
 485 490 495  
 Thr Gly Lys Ala Asn Lys Ile Thr Ile Thr Asn Asp Lys Gly Arg Leu  
 500 505 510  
 Ser Lys Glu Glu Val Glu Arg Met Val His Glu Ala Glu Gln Tyr Lys  
 515 520 525  
 Ala Glu Asp Glu Ala Gln Arg Asp Arg Val Ala Ala Lys Asn Ser Leu  
 530 535 540  
 Glu Ala His Val Phe His Val Lys Gly Ser Leu Gln Glu Glu Ser Leu  
 545 550 555 560  
 Arg Asp Lys Ile Pro Glu Glu Asp Arg Arg Lys Met Gln Asp Lys Cys  
 565 570 575  
 Arg Glu Val Leu Ala Trp Leu Glu His Asn Gln Leu Ala Glu Lys Glu  
 580 585 590  
 Glu Tyr Glu His Gln Lys Arg Glu Leu Glu Gln Ile Cys Arg Pro Ile  
 595 600 605  
 Phe Ser Arg Leu Tyr Gly Gly Pro Gly Val Pro Gly Gly Ser Ser Cys  
 610 615 620



Gly Thr Gln Ala Arg Gln Gly Asp Pro Ser Thr Gly Pro Ile Ile Glu  
 625 630 635 640

Glu Val Asp

<210> 9  
 <211> 968  
 <212> DNA  
 <213> Homo sapiens

<400> 9  
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 gcgggggtcgg ggaggtgcaa aaggatgaaa agcccgtgga cggagctgag cagatccggc 120  
 cgggctggcg gcagagaaac cgcagggaga gcctcactgc tgagcgcccc tcgacgagg 180  
 cggcagcagc ctccgtggcc tccagcatcc gacaagaagc ttcagccatg caggccccac 240  
 gggagctcgc ggtgggcatc gacctaggca ccactactc gtgctggggc gtctttcagc 300  
 agggacgctg ggagatccta gccaacgacc aaggcaaccg caccacgccc agctacgtgg 360  
 ccttcaccga caccgagcgg ctgggtcgggg acgcggccaa gaaccaggcg gccctgaacc 420  
 cccacaacac cgtgttcgat gccaaagcggc tgatcggggc caagtctcgc gacaccacgg 480  
 tgcagtcgga tatgaagcac tggcccttca aggtggtgag cggaggcgggc aagcccaagg 540  
 tgcgcgtatg ctaccgaggg gaggacaaga cgttctaccc cgaggagatc tcgtccatgg 600  
 tgctgaccaa gatgaaggag acggccgagg cgtaccttgg ccagcccgtg aagcacgcag 660  
 tgatcaccgt gccacacctat ttcagtaact cgcagcgcca agccaccaag gacgcggggg 720  
 ccatcgcggg gctcaagggtg ctgccgatca tcaatgaggc cacggcagca gccatcgctt 780  
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 gcaccttcga tgtgtcggtt ctcaccattg acgccggtgt ctttgagggtg aaagccactg 900  
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 aagaattc 968

<210> 10  
 <211> 223  
 <212> PRT  
 <213> Haemophilus ducreyi

<400> 10  
 Met Lys Lys Phe Leu Pro Ser Leu Leu Leu Met Gly Ser Val Ala Cys  
 1 5 10 15  
 Ser Ser Asn Gln Arg Met Asn Asp Tyr Ser Gln Pro Glu Ser Gln Ser  
 20 25 30  
 Asp Leu Ala Pro Lys Ser Ser Thr Ile Gln Pro Gln Pro Gln Pro Leu  
 35 40 45  
 Leu Ser Lys Thr Pro Ser Met Ser Leu Asn Leu Leu Ser Ser Ser Gly  
 50 55 60  
 Pro Asn Arg Gln Val Leu Pro Ser Glu Pro Ser Asn Phe Met Thr Leu  
 65 70 75 80  
 Met Gly Gln Asn Gly Ala Leu Leu Thr Val Trp Ala Leu Ala Lys Arg  
 85 90 95  
 Asn Trp Leu Trp Ala Tyr Pro Asn Ile Tyr Ser Gln Asp Phe Gly Asn  
 100 105 110

Ile Arg Asn Trp Lys Met Glu Pro Gly Lys His Arg Glu Tyr Phe Arg  
 115 120 125  
 Phe Val Asn Gln Ser Leu Gly Thr Cys Val Glu Ala Tyr Gly Asn Gly  
 130 135 140  
 Leu Ile His Asp Ile Cys Ser Leu Asp Lys Leu Ala Gln Glu Phe Glu  
 145 150 155 160  
 Leu Leu Pro Thr Asp Ser Gly Ala Val Val Ile Lys Ser Val Ser Gln  
 165 170 175  
 Gly Arg Cys Val Thr Tyr Asn Pro Val Ser Thr Thr Phe Tyr Ser Thr  
 180 185 190  
 Val Thr Leu Ser Val Cys Asp Gly Ala Thr Glu Pro Ser Arg Asp Gln  
 195 200 205  
 Thr Trp Tyr Leu Ala Pro Pro Val Leu Glu Ala Thr Ala Val Asn  
 210 215 220

<210> 11  
 <211> 283  
 <212> PRT  
 <213> Haemophilus ducreyi

<400> 11  
 Met Gln Trp Val Lys Gln Leu Ser Val Val Phe Cys Val Met Leu Phe  
 1 5 10 15  
 Ser Phe Ser Ser Tyr Ala Asn Leu Ser Asp Phe Lys Val Ala Thr Trp  
 20 25 30  
 Asn Leu Gln Gly Ser Ser Ala Val Asn Glu Ser Lys Trp Asn Ile Asn  
 35 40 45  
 Val Arg Gln Leu Leu Ser Gly Glu Gln Gly Ala Asp Ile Leu Met Val  
 50 55 60  
 Gln Glu Ala Gly Ser Leu Pro Ser Ser Ala Val Arg Thr Ser Arg Val  
 65 70 75 80  
 Ile Gln His Gly Gly Thr Pro Ile Glu Glu Tyr Thr Trp Asn Leu Gly  
 85 90 95  
 Thr Arg Ser Arg Pro Asn Met Val Tyr Ile Tyr Tyr Ser Arg Leu Asp  
 100 105 110  
 Val Gly Ala Asn Arg Val Asn Leu Ala Ile Val Ser Arg Arg Gln Ala  
 115 120 125  
 Asp Glu Ala Phe Ile Val His Ser Asp Ser Ser Val Leu Gln Ser Arg  
 130 135 140  
 Pro Ala Val Gly Ile Arg Ile Gly Thr Asp Val Phe Phe Thr Val His  
 145 150 155 160

Ala Leu Ala Thr Gly Gly Ser Asp Ala Val Ser Leu Ile Arg Asn Ile  
 165 170 175

Phe Thr Thr Phe Asn Ser Ser Ser Ser Pro Pro Glu Arg Arg Val Tyr  
 180 185 190

Ser Trp Met Val Val Gly Asp Phe Asn Arg Ala Pro Ala Asn Leu Glu  
 195 200 205

Val Ala Leu Arg Gln Glu Pro Ala Val Ser Glu Asn Thr Ile Ile Ile  
 210 215 220

Ala Pro Thr Glu Pro Thr His Arg Ser Gly Asn Ile Leu Asp Tyr Ala  
 225 230 235 240

Ile Leu His Asp Ala His Leu Pro Arg Arg Glu Gln Ala Arg Glu Arg  
 245 250 255

Ile Gly Ala Ser Leu Met Leu Asn Gln Leu Arg Ser Gln Ile Thr Ser  
 260 265 270

Asp His Phe Pro Val Ser Phe Val Arg Asp Arg  
 275 280

<210> 12  
 <211> 186  
 <212> PRT  
 <213> Haemophilus ducreyi

<400> 12  
 Met Lys Lys Tyr Leu Leu Ser Phe Leu Leu Ile Met Ile Leu Ala Leu  
 1 5 10 15

Ala Ser His Ala Glu Ser Asn Pro Asp Pro Thr Thr Tyr Pro Asp Val  
 20 25 30

Glu Leu Ser Pro Pro Pro Arg Ile Ser Leu Arg Ser Leu Leu Thr Ala  
 35 40 45

Gln Pro Val Lys Asn Asp His Tyr Asp Ser His Asn Tyr Leu Ser Thr  
 50 55 60

His Trp Glu Leu Ile Asp Tyr Lys Gly Lys Glu Tyr Glu Lys Leu Arg  
 65 70 75 80

Asp Gly Gly Thr Leu Val Gln Phe Lys Val Val Gly Ala Ala Lys Cys  
 85 90 95

Phe Ala Phe Leu Gly Lys Gly Thr Thr Asp Cys Lys Asp Thr Asp His  
 100 105 110

Thr Val Phe Asn Leu Ile Pro Thr Asn Thr Gly Ala Phe Leu Ile Lys  
 115 120 125

Asp Ala Leu Leu Gly Phe Cys Ile Thr Ser His Asp Phe Asp Asp Leu  
 130 135 140

Lys Leu Glu Pro Cys Gly Gly Ser Val Ser Gly Arg Thr Phe Ser Leu  
145 150 155 160

Ala Tyr Gln Trp Gly Ile Leu Pro Pro Phe Gly Pro Ser Lys Ile Leu  
165 170 175

Ile Pro Pro Val Arg Arg Asn Gln Gly Ser  
180 185

<210> 13

<211> 268

<212> PRT

<213> Campylobacter jejuni

<400> 13

Met Gln Lys Ile Ile Val Phe Ile Leu Cys Cys Phe Met Thr Phe Phe  
1 5 10 15

Leu Tyr Ala Cys Ser Ser Lys Phe Glu Asn Val Asn Pro Leu Gly Arg  
20 25 30

Ser Phe Gly Glu Phe Glu Asp Thr Asp Pro Leu Lys Leu Gly Leu Glu  
35 40 45

Pro Thr Phe Pro Thr Asn Gln Glu Ile Pro Ser Leu Ile Ser Gly Ala  
50 55 60

Asp Leu Val Pro Ile Thr Pro Ile Thr Pro Pro Leu Thr Arg Thr Ser  
65 70 75 80

Asn Ser Ala Asn Asn Asn Ala Ala Asn Gly Ile Asn Pro Arg Phe Lys  
85 90 95

Asp Glu Ala Phe Asn Asp Val Leu Ile Phe Glu Asn Arg Pro Ala Val  
100 105 110

Ser Asp Phe Leu Thr Ile Leu Gly Pro Ser Gly Ala Ala Leu Thr Val  
115 120 125

Trp Ala Leu Ala Gln Gly Asn Trp Ile Trp Gly Tyr Thr Leu Ile Asp  
130 135 140

Ser Lys Gly Phe Gly Asp Ala Arg Val Trp Gln Leu Leu Leu Tyr Pro  
145 150 155 160

Asn Asp Phe Ala Met Ile Lys Asn Ala Lys Thr Asn Thr Cys Leu Asn  
165 170 175

Ala Tyr Gly Asn Gly Ile Val His Tyr Pro Cys Asp Ala Ser Asn His  
180 185 190

Ala Gln Met Trp Lys Leu Ile Pro Met Ser Asn Thr Ala Val Gln Ile  
195 200 205

Lys Asn Leu Gly Asn Gly Lys Cys Ile Gln Ala Pro Ile Thr Asn Leu  
210 215 220

Tyr Gly Asp Phe His Lys Val Phe Lys Ile Phe Thr Val Glu Cys Ala  
 225 230 235 240

Lys Lys Asp Asn Phe Asp Gln Gln Trp Phe Leu Thr Thr Pro Pro Phe  
 245 250 255

Thr Ala Lys Pro Leu Tyr Arg Gln Gly Glu Val Arg  
 260 265

<210> 14

<211> 265

<212> PRT

<213> Campylobacter jejuni

<400> 14

Met Lys Lys Ile Ile Cys Leu Phe Leu Ser Phe Asn Leu Ala Phe Ala  
 1 5 10 15

Asn Leu Glu Asn Phe Asn Val Gly Thr Trp Asn Leu Gln Gly Ser Ser  
 20 25 30

Ala Ala Thr Glu Ser Lys Trp Ser Val Ser Val Arg Gln Leu Val Ser  
 35 40 45

Gly Ala Asn Pro Leu Asp Ile Leu Met Ile Gln Glu Ala Gly Thr Leu  
 50 55 60

Pro Arg Thr Ala Thr Pro Thr Gly Arg His Val Gln Gln Gly Gly Thr  
 65 70 75 80

Pro Ile Asp Glu Tyr Glu Trp Asn Leu Gly Thr Leu Ser Arg Pro Asp  
 85 90 95

Arg Val Phe Ile Tyr Tyr Ser Arg Val Asp Val Gly Ala Asn Arg Val  
 100 105 110

Asn Leu Ala Ile Val Ser Arg Met Gln Ala Glu Glu Val Ile Val Leu  
 115 120 125

Pro Pro Pro Thr Thr Val Ser Arg Pro Ile Ile Gly Ile Arg Asn Gly  
 130 135 140

Asn Asp Ala Phe Phe Asn Ile His Ala Leu Ala Asn Gly Gly Thr Asp  
 145 150 155 160

Val Gly Ala Ile Ile Thr Ala Val Asp Ala His Phe Ala Asn Met Pro  
 165 170 175

Gln Val Asn Trp Met Ile Ala Gly Asp Phe Asn Arg Asp Pro Ser Thr  
 180 185 190

Ile Thr Ser Thr Val Asp Arg Glu Leu Ala Asn Arg Ile Arg Val Val  
 195 200 205

Phe Pro Thr Ser Ala Thr Gln Ala Ser Gly Gly Thr Leu Asp Tyr Ala  
 210 215 220

Ile Thr Gly Asn Ser Asn Arg Gln Gln Thr Tyr Thr Pro Pro Leu Leu  
 225 230 235 240

Ala Ala Ile Leu Met Leu Ala Ser Leu Arg Ser His Ile Val Ser Asp  
 245 250 255

His Phe Pro Val Asn Phe Arg Lys Phe  
 260 265

<210> 15

<211> 189

<212> PRT

<213> Campylobacter jejuni

<400> 15

Met Lys Lys Ile Ile Thr Leu Phe Phe Met Phe Ile Thr Leu Ala Phe  
 1 5 10 15

Ala Thr Pro Thr Gly Asp Leu Lys Asp Phe Thr Glu Met Val Ser Ile  
 20 25 30

Arg Ser Leu Glu Thr Gly Ile Phe Leu Ser Ala Phe Arg Asp Thr Ser  
 35 40 45

Lys Asp Pro Ile Asp Gln Asn Trp Asn Ile Lys Glu Ile Val Leu Ser  
 50 55 60

Asp Glu Leu Lys Gln Lys Asp Lys Leu Ala Asp Glu Leu Pro Phe Gly  
 65 70 75 80

Tyr Val Gln Phe Thr Asn Pro Lys Glu Ser Asp Leu Cys Leu Ala Ile  
 85 90 95

Leu Glu Asp Gly Thr Phe Gly Ala Lys Ser Cys Gln Asp Asp Leu Lys  
 100 105 110

Asp Gly Lys Leu Glu Thr Val Phe Ser Ile Met Pro Thr Thr Thr Ser  
 115 120 125

Ala Val Gln Ile Arg Ser Leu Val Leu Glu Ser Asp Glu Cys Ile Val  
 130 135 140

Thr Phe Phe Asn Pro Asn Ile Pro Ile Gln Lys Arg Phe Gly Ile Ala  
 145 150 155 160

Pro Cys Thr Leu Asp Pro Ile Phe Phe Ala Glu Val Asn Glu Leu Met  
 165 170 175

Ile Ile Thr Pro Pro Leu Thr Ala Ala Thr Pro Leu Glu  
 180 185

<210> 16

<211> 33

<212> PRT

<213> Campylobacter jejuni

&lt;400&gt; 16

Met Leu Thr Trp Gln Gln Ile Tyr Asp Pro Phe Ser Asn Ile Trp Leu  
 1 5 10 15

Ser Ala Leu Val Ala Phe Leu Pro Ile Leu Cys Phe Leu Val Cys Leu  
 20 25 30

Val

&lt;210&gt; 17

&lt;211&gt; 237

&lt;212&gt; PRT

&lt;213&gt; Escherichia coli

&lt;400&gt; 17

Met Asp Lys Lys Leu Ile Ala Phe Leu Cys Thr Leu Ile Ile Thr Gly  
 1 5 10 15

Cys Ser Asn Gly Ile Gly Asp Ser Pro Ser Pro Pro Gly Lys Asn Val  
 20 25 30

Glu Leu Val Gly Ile Pro Gly Gln Gly Ile Ala Val Thr Ser Asn Gly  
 35 40 45

Ala Thr Pro Thr Leu Gly Ala Asn Asn Thr Glu Phe Pro Glu Val Ser  
 50 55 60

Ile Met Ser Thr Gly Gly Ala Leu Leu Thr Ile Trp Ala Arg Pro Val  
 65 70 75 80

Arg Asn Trp Leu Trp Gly Tyr Thr Pro Phe Asp Ser Val Asn Phe Gly  
 85 90 95

Glu Asn Arg Asn Trp Lys Val Val Asp Gly Lys Asp Ala Gly Thr Val  
 100 105 110

Lys Phe Val Asn Val Ala Gln Gly Thr Cys Met Glu Ala Phe Lys Asn  
 115 120 125

Gly Val Ile His Asn Thr Cys Asp Asp Asn Ser Leu Ser Gln Glu Phe  
 130 135 140

Gln Leu Leu Pro Ser Thr Asn Gly Asn Val Leu Ile Arg Ser Ser Ala  
 145 150 155 160

Leu Gln Thr Cys Ile Arg Ala Asp Tyr Leu Ser Arg Thr Ile Leu Ser  
 165 170 175

Pro Phe Ala Phe Thr Ile Thr Leu Glu Lys Cys Pro Gly Ala Lys Glu  
 180 185 190

Glu Thr Gln Glu Met Leu Trp Ala Ile Ser Pro Pro Val Arg Ala Ala  
 195 200 205

Lys Pro Asn Leu Ile Lys Pro Glu Leu Arg Pro Phe Arg Pro Leu Pro  
 210 215 220

Ile Pro Pro His Asp Lys Pro Asp Gly Met Glu Gly Val  
 225 230 235

<210> 18

<211> 273

<212> PRT

<213> Escherichia coli

<400> 18

Met Lys Lys Leu Leu Phe Leu Leu Met Ile Leu Pro Gly Ile Ser Phe  
 1 5 10 15

Ala Asp Leu Ser Asp Phe Lys Val Ala Thr Trp Asn Leu Gln Gly Ser  
 20 25 30

Asn Ala Pro Thr Glu Asn Lys Trp Asn Thr His Val Arg Gln Leu Val  
 35 40 45

Thr Gly Ser Gly Ala Val Asp Ile Leu Met Val Gln Glu Ala Gly Ala  
 50 55 60

Val Pro Ala Ser Ala Thr Leu Thr Glu Arg Glu Phe Ser Thr Pro Gly  
 65 70 75 80

Ile Pro Met Asn Glu Tyr Ile Trp Asn Thr Gly Thr Asn Ser Arg Pro  
 85 90 95

Gln Glu Leu Phe Ile Tyr Phe Ser Arg Val Asp Ala Phe Ala Asn Arg  
 100 105 110

Val Asn Leu Ala Ile Val Ser Asn Arg Arg Ala Asp Glu Val Ile Val  
 115 120 125

Leu Pro Pro Pro Thr Val Val Ser Arg Pro Ile Ile Gly Ile Arg Ile  
 130 135 140

Gly Asn Asp Val Phe Phe Ser Thr His Ala Leu Ala Asn Arg Gly Val  
 145 150 155 160

Asp Ser Gly Ala Ile Val Asn Ser Val Phe Glu Phe Phe Asn Arg Gln  
 165 170 175

Thr Asp Pro Ile Arg Gln Ala Ala Asn Trp Met Ile Ala Gly Asp Phe  
 180 185 190

Asn Arg Ser Pro Ala Thr Leu Phe Ser Thr Leu Glu Pro Gly Ile Arg  
 195 200 205

Asn His Val Asn Ile Ile Ala Pro Pro Asp Pro Thr Gln Ala Ser Gly  
 210 215 220

Gly Val Leu Asp Tyr Ala Val Val Gly Asn Ser Val Ser Phe Val Leu  
 225 230 235 240



Pro Leu Leu Arg Ala Ser Leu Leu Phe Gly Leu Leu Arg Gly Gln Ile  
                                   245                                  250                                  255

Ala Ser Asp His Phe Pro Val Gly Phe Ile Pro Gly Arg Gly Ala Arg  
                                   260                                  265                                  270

Arg

<210> 19

<211> 190

<212> PRT

<213> Escherichia coli

<400> 19

Met Lys Thr Val Ile Val Phe Phe Val Leu Leu Leu Thr Gly Cys Ala  
   1                                  5                                  10                                  15

Ser Glu Pro Ala Asn Gln Arg Asn Leu Leu Thr Gln Phe Val Gly Asn  
                                   20                                  25                                  30

Asn Ala Pro Val Asp Pro Glu Pro Ser Pro Val Leu Val Asn Ile Arg  
                                   35                                  40                                  45

Asn Val Leu Thr Gly Gly Ile Ile Arg Asn Pro Val Gly Ser Asp Phe  
   50                                  55                                  60

Asn Val Asn Asn Trp Val Ile Ser Glu Val Lys Thr Asn Asp Leu Asp  
   65                                  70                                  75                                  80

Leu Ile Ser Ala Pro Gly Gly His Val Gln Ile Lys Asn Pro Asp Gly  
                                   85                                  90                                  95

Asn Glu Cys Phe Ala Ile Leu Asn Gly Gln Leu Ala Val Ala Lys Gln  
                                   100                                  105                                  110

Cys Ser Glu Ser Asp Arg Asn Ala Leu Phe Thr Phe Ile Thr Ser Asp  
                                   115                                  120                                  125

Thr Gly Ala Val Gln Ile Lys Ser Ile Gly Ser Gly Gln Cys Leu Gly  
   130                                  135                                  140

Asn Gly Glu Ser Ile Thr Asp Phe Arg Leu Lys Lys Cys Val Asp Asp  
   145                                  150                                  155                                  160

Leu Gly Arg Pro Phe Asp Thr Val Pro Pro Gly Leu Leu Trp Met Leu  
                                   165                                  170                                  175

Asn Pro Pro Leu Ser Pro Ala Ile Met Ser Pro Leu Thr Ser  
                                   180                                  185                                  190

<210> 20

<211> 258

<212> PRT

<213> Escherichia coli

&lt;400&gt; 20

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Met Ala Asn Lys Arg Thr Pro Ile Phe Ile Ala Gly Ile Leu Ile Pro
 1           5           10           15

Ile Leu Leu Asn Gly Cys Ser Ser Gly Lys Asn Lys Ala Tyr Leu Asp
          20           25           30

Pro Lys Val Phe Pro Pro Gln Val Glu Gly Gly Pro Thr Val Pro Ser
          35           40           45

Pro Asp Glu Pro Gly Leu Pro Leu Pro Gly Pro Gly Pro Ala Leu Pro
          50           55           60

Thr Asn Gly Ala Ile Pro Ile Pro Glu Pro Gly Thr Ala Pro Ala Val
          65           70           75           80

Ser Leu Met Asn Met Asp Gly Ser Val Leu Thr Met Trp Ser Arg Gly
          85           90           95

Ala Gly Ser Ser Leu Trp Ala Tyr Tyr Ile Gly Asp Ser Asn Ser Phe
          100          105          110

Gly Glu Leu Arg Asn Trp Gln Ile Met Pro Gly Thr Arg Pro Asn Thr
          115          120          125

Ile Gln Phe Arg Asn Val Asp Val Gly Thr Cys Met Thr Ser Phe Pro
          130          135          140

Gly Phe Lys Gly Gly Val Gln Leu Ser Thr Ala Pro Cys Lys Phe Gly
          145          150          155          160

Pro Glu Arg Phe Asp Phe Gln Pro Met Ala Thr Arg Asn Gly Asn Tyr
          165          170          175

Gln Leu Lys Ser Leu Ser Thr Gly Leu Cys Ile Arg Ala Asn Phe Leu
          180          185          190

Gly Arg Thr Pro Ser Ser Pro Tyr Ala Thr Thr Leu Thr Met Glu Arg
          195          200          205

Cys Pro Ser Ser Gly Glu Lys Asn Phe Glu Phe Met Trp Ser Ile Ser
          210          215          220

Glu Pro Leu Arg Pro Ala Leu Ala Thr Ile Ala Lys Pro Glu Ile Arg
          225          230          235          240

Pro Phe Pro Pro Gln Pro Ile Glu Pro Asp Glu His Ser Thr Gly Gly
          245          250          255

Glu Gln

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&lt;210&gt; 21

&lt;211&gt; 269

&lt;212&gt; PRT

&lt;213&gt; Escherichia coli

&lt;400&gt; 21

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Met Lys Lys Tyr Ile Ile Ser Leu Ile Val Phe Leu Ser Phe Tyr Ala
 1           5           10           15

Gln Ala Asp Leu Thr Asp Phe Arg Val Ala Thr Trp Asn Leu Gln Gly
          20           25           30

Ala Ser Ala Thr Thr Glu Ser Lys Trp Asn Ile Asn Val Arg Gln Leu
          35           40           45

Ile Ser Gly Glu Asn Ala Val Asp Ile Leu Ala Val Gln Glu Ala Gly
          50           55           60

Ser Pro Pro Ser Thr Ala Val Asp Thr Gly Thr Leu Ile Pro Ser Pro
          65           70           75           80

Gly Ile Pro Val Arg Glu Leu Ile Trp Asn Leu Ser Thr Asn Ser Arg
          85           90           95

Pro Gln Gln Val Tyr Ile Tyr Phe Ser Ala Val Asp Ala Leu Gly Gly
          100          105          110

Arg Val Asn Leu Ala Leu Val Ser Asn Arg Arg Ala Asp Glu Val Phe
          115          120          125

Val Leu Ser Pro Val Arg Gln Gly Gly Arg Pro Leu Leu Gly Ile Arg
          130          135          140

Ile Gly Asn Asp Ala Phe Phe Thr Ala His Ala Ile Ala Met Arg Asn
          145          150          155          160

Asn Asp Ala Pro Ala Leu Val Glu Glu Val Tyr Asn Phe Phe Arg Asp
          165          170          175

Ser Arg Asp Pro Val His Gln Ala Leu Asn Trp Met Ile Leu Gly Asp
          180          185          190

Phe Asn Arg Glu Pro Ala Asp Leu Glu Met Asn Leu Thr Val Pro Val
          195          200          205

Arg Arg Ala Ser Glu Ile Ile Ser Pro Ala Ala Ala Thr Gln Thr Ser
          210          215          220

Gln Arg Thr Leu Asp Tyr Ala Val Ala Gly Asn Ser Val Ala Phe Arg
          225          230          235          240

Pro Ser Pro Leu Gln Ala Gly Ile Val Tyr Gly Ala Arg Arg Thr Gln
          245          250          255

Ile Ser Ser Asp His Phe Pro Val Gly Val Ser Arg Arg
          260          265

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&lt;210&gt; 22

&lt;211&gt; 181

&lt;212&gt; PRT

&lt;213&gt; Escherichia coli

&lt;400&gt; 22

Met Lys Lys Leu Ala Ile Val Phe Thr Met Leu Leu Ile Ala Gly Cys  
 1 5 10 15

Ser Ser Ser Gln Asp Ser Ala Asn Asn Gln Ile Asp Glu Leu Gly Lys  
 20 25 30

Glu Asn Asn Ser Leu Phe Thr Phe Arg Asn Ile Gln Ser Gly Leu Met  
 35 40 45

Ile His Asn Gly Leu His Gln His Gly Arg Glu Thr Ile Gly Trp Glu  
 50 55 60

Ile Val Pro Val Lys Thr Pro Glu Glu Ala Leu Val Thr Asp Gln Ser  
 65 70 75 80

Gly Trp Ile Met Ile Arg Thr Pro Asn Thr Asp Gln Cys Leu Gly Thr  
 85 90 95

Pro Asp Gly Arg Asn Leu Leu Lys Met Thr Cys Asn Ser Thr Ala Lys  
 100 105 110

Lys Thr Leu Phe Ser Leu Ile Pro Ser Thr Thr Gly Ala Val Gln Ile  
 115 120 125

Lys Ser Val Leu Ser Gly Leu Cys Phe Leu Asp Ser Lys Asn Ser Gly  
 130 135 140

Leu Ser Phe Glu Thr Gly Lys Cys Ile Ala Asp Phe Lys Lys Pro Phe  
 145 150 155 160

Glu Val Val Pro Gln Ser His Leu Trp Met Leu Asn Pro Leu Asn Thr  
 165 170 175

Glu Ser Pro Ile Ile  
 180

&lt;210&gt; 23

&lt;211&gt; 155

&lt;212&gt; PRT

&lt;213&gt; Escherichia coli

&lt;400&gt; 23

Glu Asn Lys Trp Asn Thr His Val Arg Gln Leu Val Thr Gly Ser Gly  
 1 5 10 15

Ala Val Asp Ile Leu Met Val Gln Glu Ala Gly Ala Val Pro Ala Ser  
 20 25 30

Ala Thr Leu Thr Glu Arg Glu Phe Ser Thr Pro Gly Ile Pro Met Asn  
 35 40 45

Glu Tyr Ile Trp Asn Thr Gly Thr Asn Ser Arg Pro Gln Glu Leu Phe  
 50 55 60

Ile Tyr Phe Ser Arg Val Asp Ala Phe Ala Asn Arg Val Asn Leu Ala  
 65 70 75 80

Ile Val Ser Asn Arg Arg Ala Asp Glu Val Ile Val Leu Pro Pro Pro  
85 90 95

Thr Val Val Ser Arg Pro Ile Ile Gly Ile Arg Ile Gly Asn Asp Val  
100 105 110

Phe Phe Ser Thr His Ala Leu Ala Asn Arg Gly Val Asp Ser Gly Ala  
115 120 125

Ile Val Asn Ser Val Phe Glu Phe Phe Asn Arg Gln Thr Asp Pro Ile  
130 135 140

Arg Gln Ala Ala Asn Trp Met Ile Ala Gly Asp  
145 150 155

<210> 24  
<211> 609  
<212> PRT  
<213> Homo sapiens

<400> 24  
Met Ser Gly Trp Glu Ser Tyr Tyr Lys Thr Glu Gly Asp Glu Glu Ala  
1 5 10 15

Glu Glu Glu Gln Glu Glu Asn Leu Glu Ala Ser Gly Asp Tyr Lys Tyr  
20 25 30

Ser Gly Arg Asp Ser Leu Ile Phe Leu Val Asp Ala Ser Lys Ala Met  
35 40 45

Phe Glu Ser Gln Ser Glu Asp Glu Leu Thr Pro Phe Asp Met Ser Ile  
50 55 60

Gln Cys Ile Gln Ser Val Tyr Ile Ser Lys Ile Ile Ser Ser Asp Arg  
65 70 75 80

Asp Leu Leu Ala Val Val Phe Tyr Gly Thr Glu Lys Asp Lys Asn Ser  
85 90 95

Val Asn Phe Lys Asn Ile Tyr Val Leu Gln Glu Leu Asp Asn Pro Gly  
100 105 110

Ala Lys Arg Ile Leu Glu Leu Asp Gln Phe Lys Gly Gln Gln Gly Gln  
115 120 125

Lys Arg Phe Gln Asp Met Met Gly His Gly Ser Asp Tyr Ser Leu Ser  
130 135 140

Glu Val Leu Trp Val Cys Ala Asn Leu Phe Ser Asp Val Gln Phe Lys  
145 150 155 160

Met Ser His Lys Arg Ile Met Leu Phe Thr Asn Glu Asp Asn Pro His  
165 170 175

Gly Asn Asp Ser Ala Lys Ala Ser Arg Ala Arg Thr Lys Ala Gly Asp  
180 185 190

Leu Arg Asp Thr Gly Ile Phe Leu Asp Leu Met His Leu Lys Lys Pro  
 195 200 205  
 Gly Gly Phe Asp Ile Ser Leu Phe Tyr Arg Asp Ile Ile Ser Ile Ala  
 210 215 220  
 Glu Asp Glu Asp Leu Arg Val His Phe Glu Glu Ser Ser Lys Leu Glu  
 225 230 235 240  
 Asp Leu Leu Arg Lys Val Arg Ala Lys Glu Thr Arg Lys Arg Ala Leu  
 245 250 255  
 Ser Arg Leu Lys Leu Lys Leu Asn Lys Asp Ile Val Ile Ser Val Gly  
 260 265 270  
 Ile Tyr Asn Leu Val Gln Lys Ala Leu Lys Pro Pro Pro Ile Lys Leu  
 275 280 285  
 Tyr Arg Glu Thr Asn Glu Pro Val Lys Thr Lys Thr Arg Thr Phe Asn  
 290 295 300  
 Thr Ser Thr Gly Gly Leu Leu Leu Pro Ser Asp Thr Lys Arg Ser Gln  
 305 310 315 320  
 Ile Tyr Gly Ser Arg Gln Ile Ile Leu Glu Lys Glu Glu Thr Glu Glu  
 325 330 335  
 Leu Lys Arg Phe Asp Asp Pro Gly Leu Met Leu Met Gly Phe Lys Pro  
 340 345 350  
 Leu Val Leu Leu Lys Lys His His Tyr Leu Arg Pro Ser Leu Phe Val  
 355 360 365  
 Tyr Pro Glu Glu Ser Leu Val Ile Gly Ser Ser Thr Leu Phe Ser Ala  
 370 375 380  
 Leu Leu Ile Lys Cys Leu Glu Lys Glu Val Ala Ala Leu Cys Arg Tyr  
 385 390 395 400  
 Thr Pro Arg Arg Asn Ile Pro Pro Tyr Phe Val Ala Leu Val Pro Gln  
 405 410 415  
 Glu Glu Glu Leu Asp Asp Gln Lys Ile Gln Val Thr Pro Pro Gly Phe  
 420 425 430  
 Gln Leu Val Phe Leu Pro Phe Ala Asp Asp Lys Arg Lys Met Pro Phe  
 435 440 445  
 Thr Glu Lys Ile Met Ala Thr Pro Glu Gln Val Gly Lys Met Lys Ala  
 450 455 460  
 Ile Val Glu Lys Leu Arg Phe Thr Tyr Arg Ser Asp Ser Phe Glu Asn  
 465 470 475 480  
 Pro Val Leu Gln Gln His Phe Arg Asn Leu Glu Ala Leu Ala Leu Asp  
 485 490 495

Leu	Met	Glu	Pro	Glu	Gln	Ala	Val	Asp	Leu	Thr	Leu	Pro	Lys	Val	Glu
			500					505					510		
Ala	Met	Asn	Lys	Arg	Leu	Gly	Ser	Leu	Val	Asp	Glu	Phe	Lys	Glu	Leu
		515					520					525			
Val	Tyr	Pro	Pro	Asp	Tyr	Asn	Pro	Glu	Gly	Lys	Val	Thr	Lys	Arg	Lys
	530					535					540				
His	Asp	Asn	Glu	Gly	Ser	Gly	Ser	Lys	Arg	Pro	Lys	Val	Glu	Tyr	Ser
545					550					555					560
Glu	Glu	Glu	Leu	Lys	Thr	His	Ile	Ser	Lys	Gly	Thr	Leu	Gly	Lys	Phe
				565					570					575	
Thr	Val	Pro	Met	Leu	Lys	Glu	Ala	Cys	Arg	Ala	Tyr	Gly	Leu	Lys	Ser
			580					585					590		
Gly	Leu	Lys	Lys	Gln	Glu	Leu	Leu	Glu	Ala	Leu	Thr	Lys	His	Phe	Gln
		595					600					605			

Asp

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<210> 25
<211> 247
<212> PRT
<213> Homo sapiens
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<400> 25																
Met	Gln	Ala	Pro	Arg	Glu	Leu	Ala	Val	Gly	Ile	Asp	Leu	Gly	Thr	Thr	
1				5					10					15		
Tyr	Ser	Cys	Val	Gly	Val	Phe	Gln	Gln	Gly	Arg	Val	Glu	Ile	Leu	Ala	
			20					25					30			
Asn	Asp	Gln	Gly	Asn	Arg	Thr	Thr	Pro	Ser	Tyr	Val	Ala	Phe	Thr	Asp	
		35					40					45				
Thr	Glu	Arg	Leu	Val	Gly	Asp	Ala	Ala	Lys	Asn	Gln	Ala	Ala	Leu	Asn	
	50					55					60					
Pro	His	Asn	Thr	Val	Phe	Asp	Ala	Lys	Arg	Leu	Ile	Gly	Arg	Lys	Phe	
65					70					75					80	
Ala	Asp	Thr	Thr	Val	Gln	Ser	Asp	Met	Lys	His	Trp	Pro	Phe	Lys	Val	
				85					90					95		
Val	Ser	Gly	Gly	Gly	Lys	Pro	Lys	Val	Arg	Val	Cys	Tyr	Arg	Gly	Glu	
			100					105					110			
Asp	Lys	Thr	Phe	Tyr	Pro	Glu	Glu	Ile	Ser	Ser	Met	Val	Leu	Thr	Lys	
		115					120					125				
Met	Lys	Glu	Thr	Ala	Glu	Ala	Tyr	Leu	Gly	Gln	Pro	Val	Lys	His	Ala	
	130					135					140					

